

SEQUENCE LISTING

EPICYTE PHARMACEUTICAL, INC. HIATT, ANDREW C. HEIN, MICH B. FITCHEN, JOHN H.



120> NOVEL EPITHELIAL TISSUE IMAGING AGENT

<130> 068904-0204

<140> UNASSIGNED

<141> HEREWITH

<150> 09/005,167

<151> 1998-01-09

<150> 08/702,480

<151> 1997-01-10

<160> 93

<170> PatentIn 3.0

<210> 1

<211> 137

<212> Protein

<213> Human

<220>

<221> misc-feature

<222> Synthetic polypeptide J chain

<400> 1

Gln Glu Asp Glu Arg Ile Val Leu Val Asp Asn Lys Cys Lys Cys Ala 1 5 10 15

Arg Ile Thr Ser Arg Ile Ile Arg Ser Ser Glu Asp Pro Asn Glu Asp 20 25 30

Ilé Val Glu Arg Asn Ile Arg Ile Ile Val Pro Leu Asn Asn Arg Glu 35 40 45

Asn Ile Ser Asp Pro Thr Ser Pro Leu Arg Thr Arg Pro Val Tyr His 50 55

Leu Ser Asp Leu Cys Lys Lys Cys Asp Pro Thr Glu Val Glu Leu Asp 65 70 75 80

Asn Gln Ile Val Thr Ala Thr Gln Ser Asn Ile Cys Asp Glu Asp Ser 85 90 95 . 3.

Ala Thr Glu Thr Cys Tyr Thr Tyr Asp Arg Asn Lys Cys Tyr Thr Ala 100 105 110

Val Val Pro Leu Val Tyr Gly Gly Glu Thr Lys Met Val Glu Thr Ala 115 120 125

Leu Thr Pro Asp Ala Cys Tyr Pro Asp 130 135

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Arg Ile Thr Ser Arg Ile Ile Pro Ser Ala Glu Asp Pro Ser Gln Asp 20 25 30

Ile Val Glu Arg Asn Val Arg Ile Ile Val Pro Leu Asn Ser Arg Glu
35 40 45

Asn Ile Ser Asp Pro Thr Ser Pro Met Arg Thr Lys Pro Val Tyr His 50 60

Leu Ser Asp Leu Cys Lys Lys Cys Asp Thr Thr Glu Val Glu Leu Glu 65 70 75 80

Asp Gln Val Val Thr Ala Ser Gln Ser Asn Ile Cys Asp Ser Asp Ala 85 90 95

Glu Thr Cys Tyr Thr Tyr Asp Arg Asn Lys Cys Tyr Thr Asn Arg Val

Lys Leu Ser Tyr Arg Gly Gln Thr Lys Met Val Glu Thr Ala Leu Thr 115 120 125

Pro Asp Ser Cys Tyr Pro Asp 130 135

<210> 3

<211> 137 <212> Protein

<213> Rabbit

<220>

<221> misc-feature

<222> Synthetic polypeptide J chain

<400> 3

Asp Asp Glu Ala Thr Ile Leu Ala Asp Asn Lys Cys Met Cys Thr Arg 1 5 10 15

Val Thr Ser Arg Ile Ile Pro Ser Thr Glu Asp Pro Asn Glu Asp Ile 20 25 30

Val Glu Arg Asn Ile Arg Ile Val Val Pro Leu Asn Asn Arg Glu Asn 35 40 45

Ile Ser Asp Pro Thr Ser Pro Leu Arg Arg Asn Pro Val Tyr His Leu 50 55 60

Ser Asp Val Cys Lys Lys Cys Asp Pro Val Glu Val Glu Leu Glu Asp
65 70 75 80

Gln Val Val Thr Ala Thr Gln Ser Asn Ile Cys Asn Glu Asp Asp Gly 85 90 95

Val Pro Glu Thr Cys Tyr Met Tyr Asp Arg Asn Lys Cys Tyr Thr Thr 100 105 110

Met Val Pro Leu Arg Tyr His Gly Glu Thr Lys Met Val Gln Ala Ala 115 120 125

Leu Thr Pro Asp Ser Cys Tyr Pro Asp

<210> 4
<211> 136

<212> Protein

<213> Bovine

<220>

* 52*

<221> misc-feature

<222> Synthetic polypeptide J chain

<400> 4

Glu Asp Glu Ser Thr Val Leu Val Asp Asn Lys Cys Gln Cys Val Arg 1 5 10 15

Ile Thr Ser Arg Ile Ile Arg Asp Pro Asp Asn Pro Ser Glu Asp Ile
20 25 30

Val Glu Arg Asn Ile Arg Ile Ile Val Pro Leu Asn Thr Arg Glu Asn 35 40 45

Ile Ser Asp Pro Thr Ser Pro Leu Arg Thr Glu Pro Lys Tyr Asn Leu 50 55 60

Ala Asn Leu Cys Lys Lys Cys Asp Pro Thr Glu Ile Glu'Leu Asp Asn 65 70 75 80

Gln Val Phe Thr Ala Ser Gln Ser Asn Ile Cys Pro Asp Asp Asp Tyr 95 90 95

Ser Glu Thr Cys Tyr Thr Tyr Asp Arg Asn Lys Cys Tyr Thr Thr Leu 100 105 110

Val Pro Ile Thr. His Arg Gly Val Thr Arg Met Val Lys Ala Thr Leu 115 120 125

Thr Pro Asp Ser Cys Tyr Pro Asp 130 135

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<213> Bull frog
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     Ser Ser Arg Phe Val Pro Ser Thr Glu Arg Pro Gly Glu Glu Ile Leu
                                    25
    Glu Arg Asn Ile Gln Ile Thr Ile Pro Thr Ser Ser Arg Met Xaa Ile
            35
    Ser Asp Pro Tyr Ser Pro Leu Arg Thr Gln Pro Val Tyr Asn Leu Trp
    Asp Ile Cys Gln Lys Cys Asp Pro Val Gln Leu Glu Ile Gly Gly Ile
    Pro Val Leu Ala Ser Gln Pro Xaa Xaa Ser Xaa Pro Asp Asp Glu Cys
    Tyr Thr Thr Glu Val Asn Phe Lys Lys Val Pro Leu Thr Pro Asp
                                    105
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            115
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<211> 129
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Asn Gln Pro Val Tyr His Leu Ser Pro Ser Cys Lys Lys Cys Asp Pro 50 55 60

Tyr Glu Asp Gly Val Val Thr Ala Thr Glu Thr Asm Ile Cys Tyr Pro 65 70 75 80

Asp Gln Gly Val Pro Gln Ser Cys Arg Asp Tyr Cys Pro Glu Leu Asp 85 90 95

Arg Asn Lys Cys Tyr Thr Val Leu Val Pro Pro Gly Tyr Thr Gly Glu
100 105 110

Thr Lys Met Val Gln Asn Ala Leu Thr Pro Asp Ala Cys Tyr Pro Asp 115 120 125

<210> 7
<211> 421
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<220>
<221> CDS
<222> (1)..(414)
<220>
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<222> Description of Artifit target of "full length" TM

<222> Description of Artificial Sequence: Synthetic polypeptide including target of "full length" TM cDNA

20 25 30

GAT ATA GTC GAA CGT AAC ATC CGT ATC ATC GTC CCA CTG AAT AAC CGG
Asp Ile Val Glu Arg Asn Ile Arg Ile Ile Val Pro Leu Asn Asn Arg

40

GAG AAT ATC TCA GAT CCT ACA AGT CCG TTG CGC ACA CGC TTC GTA TAC
Glu Asn Ile Ser Asp Pro Thr Ser Pro Leu Arg Thr Arg Phe Val Tyr
50 55 60

CAC CTG TCA GAT CTG TGT AAG AAG TGT GAT CCA ACA GAG GTA GAG CTG

His Leu Ser Asp Leu Cys Lys Lys Cys Asp Pro Thr Glu Val Glu Leu

65 70 75 80

Двр			ATA Ile											GAG. Glu 95		288
			GAA Glu 100													336
			CCG Pro													384
•			CCC Pro							TGA	₹TT C					421
<21	1> : 2> 1 3> ;	ANC	Eicia	al Se	equer	nce										
<22 <22	2>	CD\$ (1).	. (21	3)												
	l> 1 2> 1	-	-feat		of A	rtif	icia	l Se	ğuen	ce: l	Nucl	eoti	ie s	equei	nce o	f Core TM
<22 <22 CDN <40 GAT	1> 1 2> 1 A 0> (Desci B AAG		ion o	TGT	GCT	CGT	ATT	ACT	TCT	AGA	ATC	ATC	CGT	AGC	of Core TM
<22 <22 CDN <40 GAT Asp 1	1> T 2> 1 A CAG Gln	AAG Lys GAC	ript: TGC	AAG Lys 5	TGT Cys GAA	GCT Ala GAT	CGT Arg	ATT Ile	ACT Thr 10 GAA Glu	TCT Ser	AGA Arg	ATC Ile	ATC Ile	CGT Arg 15	AGC Ser	
<22 <22 CDN <40 GAT Asp 1 TCA Ser	l> r 2> l A CAG Gln GAG Glu	AAG Lys GAC Asp	TGC Cys CCA Pro	AAG Lys 5 AAT ASD	TGT Cys GAA Glu	GCT Ala GAT ABP	CGT Arg ATA Ile	ATT Ile GTC Val 25	ACT Thr 10 GAA Glu	TCT ser CGT Arg	AGA Arg AAC ABN	ATC Ile	ATC Ile CGT Arg 30	CGT Arg 15 ATC Ile	AGC Ser ATC Ile	48
<222 <221 CDN <40 GAT Asp 1 TCA Ser GTC Val	l> r 2> l A CAG Gln GAG Glu CCA Pro	AAG Lys GAC Asp CTG Leu 35	TGC Cys CCA Pro 20	AAG Lys 5 AAT ASN AAC ASN	TGT Cys GAA Glu CGG Arg	GCT Ala GAT ABP GAG Glu	CGT Arg ATA Ile AAT Asn 40	ATT Ile GTC Val 25 ATC Ile	ACT Thr 10 GAA Glu TCA Ser	TCT Ser CGT Arg GAT Asp	AGA Arg AAC ABN CCT Pro	ATC Ile ATC Ile ATC ACA Thr 45	ATC Ile CGT Arg 30 AGT Ser	CGT Arg 15 ATC Ile CCG Pro	AGC Ser ATC Ile TTG Leu	48 96

<210> 9 <211> 140 <212> DNA <213> Artificial Sequence	
<220> <221> misc-feature <222> Description of Artificial Sequence: Nucleotide sequence of C2	fragment
<400> 9 CTAGAATCAT CCGTAGCTCA GAGGACCCAA ATGAAGATAT AGTCGAACGT AACATCCGTA	60
TCATCGTCCC ACTGAATAAC CGGGAGAATA TCTCAGATCC TACAAGTCCG TTGCGCACAC	120
GCTTCGTATA CCACCTGTCA	140
<210> 10 <211> 31 <212> DNA <213> Artificial Sequence	
<220> <221> misc-feature <222> Description of Artificial Sequence: Nucleotide sequence of D1. fragment	1
<400> 10 GATCAGAAGT GCAAGTGTGC TCGTATTACT T	31 .
<210> 11 <211> 44 <212> DNA <213> Artificial Sequence	
<220> <221> CDS <222> (1)(42)	
<220> <221> misc-feature <222> Description of Artificial Sequence: Nucleotide sequence of L3D fragment	
<pre><400> 11 GAT CTG TGT AAG AAG GAT GAA GAT TCC GCT ACA GAA ACC TGC Asp Leu Cys Lys Lys Asp Glu Asp Ser Ala Thr Glu Thr Cys</pre>	42

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<210> 12 <211> 109 <212> DNA <213> Artificial Sequence	
<220> <221> misc-feature <222> Description of Artificial Sequence: Nucleotide sequence of T4 fe	ragment .
<400> 12 GCACCTACGA TAGGAACAAA TGCTACACGG CCGTGGTTCC GCTCGTGTAT GGTGGAGAGA	60
CAAAAATGGT GGAAACTGCC CTTACGCCCG ATGCATGCTA CCCTGACTG	109
<210> 13 <211> 286 <212> DNA <213> Artificial Sequence	
<220> <221> CDS <222> (1)(282)	
<220> <221> misc-feature <222> Description of Artificial Sequence: Nucleotide sequence of Core cDNA using L3	TM
<400> 13 GAC AAC AAG TGC AAG TGT GCT CGT ATT ACT TCT AGA ATC ATC CGT AGC	48
Asp Asn Lys Cys Lys Cys Ala Arg Ile Thr Ser Arg Ile Ile Arg Ser 15 20 25 30	
TCA GAG GAC CCA AAT GAA GAT ATA GTC GAA CGT AAC ATC CGT ATC ATC Ser Glu Asp Pro Asn Glu Asp Ile Val Glu Arg Asn Ile Arg Ile Ile 35 40 45	96
GTC CCA CTG AAT AAC CGG GAG AAT ATC TCA GAT CCT ACA AGT CCG TTG Val Pro Leu Asn Asn Arg Glu Asn Ile Ser Asp Pro Thr Ser Pro Leu 50 55 60	144
CGC ACA CGC TTC GTA TAC CAC CTG TCA GAT CTG TGT AAG AAG TGT GAT Arg Thr Arg Phe Val Tyr His Leu Ser Asp Leu Cys Lys Lys Cys Asp	192

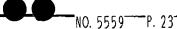
70

85

CCA ACA GAG GTA GAG CTG GAC AAT CAG ATA GTC ACT GCG ACT CAA AGC

Pro Thr Glu Val Glu Leu Asp Asn Gln Ile Val Thr Ala Thr Gln Ser

65



61

AAC ATT TGC GAT GAG GAC AGC GCT ACA GAA ACC TGC TAC TGA 282 Asn Ile Cys Asp Glu Asp Ser Ala Thr Glu Thr Cys Tyr * 95 100 ATTC 286 <210> 14 <211> 105 <212> DNA <213> Artificial Sequence <220> <221> CDS <222> (1)..(105) <220> <221> misc-feature <222> Description of Artificial Sequence: Nucleotide sequence of L3 fragment <400> 14 GAT CTG TGT AAG AAG TGT GAT CCA ACA GAG GTA GAG CTG GAC AAT CAG 48 Asp Leu Cys Lys Lys Cys Asp Pro Thr Glu Val Glu Leu Asp Asn Gln 100 105 ATA GTC ACT GCG ACT CAA AGC AAC ATT TGC GAT GAG GAC AGC GCT ACA 96 Ile Val Thr Ala Thr Gln Ser Asn Ile Cys Asp Glu Asp Ser Ala Thr 115 CTT TGG ACG 105 Leu Trp Thr <210> 15 <211> 61 <212> DNA <213> Artificial Sequence <220> <221> misc-feature <222> Description of Artificial Sequence: Nucleotide sequence of D1 fragment GATCAGGAAG ATGAACGTAT TGTTCTGGTT GACAACAAGT GCAAGTGTGC TCGTATTACT 60

*40

<210> 16 <211> 61 <212> DNA <213> Artificial Sequence	
<220> <221> misc-feature <222> Description of Artificial Sequence: Nucleotide sequence of Tp	3 2
<400> 16 GCGATGACGA CGATAAGGCC CAAACGGAGA CCTGTACTGT TGCGCCTCGT GAACGGCAAA	60
ACTGCGGATT CCCGGAAGTA ACACCCTCTC AGTGCGCTAA TAAAGGCTGC TGTTTTGATG	120
ACACGGTACG GGGCGTTCCG TGGTGCTTCT ACCCCAATAC AATTGACGTT CCGCCTGAAG	180
AAGAGTGCGA GCCGTAAG	198
<210> 17 <211> 138 <212> Protein <213> Artificial Sequence <220>	
<pre><221> misc-feature <222> Description of Artificial Sequence: Synthetic polypeptide of length" TM cDNA</pre>	"full
<400> 17	
Asp Gln Glu Asp Glu Arg Ile Val Leu Val Asp Asn Lys Cys Lys Cys 1 10 15	
Ala Arg Ile Thr Ser Arg Ile Ile Arg Ser Ser Glu Asp Pro Asn Glu 20 25 30	
Asp Ile Val Glu Arg Asn Ile Arg Ile Ile Val Pro Leu Asn Asn Arg 35 40 45	
Glu Asn Ile Ser Asp Pro Thr Ser Pro Leu Arg Thr Arg Phe Val Tyr 50 55 60 .	
His Leu Ser Asp Leu Cys Lys Lys Cys Asp Pro Thr Glu Val Glu Leu 65 75 80	

Ser Ala Thr Glu Thr Cys Ser Thr Tyr Asp Arg Asn Lys Cys Tyr Thr

Ala Val Val Pro Leu Val Tyr Gly Gly Glu Thr Lys Met Val Glu Thr 120

Ala L u Thr Pro Asp Ala Cys Tyr Pro Asp 130 135

<210> 18

<211> 71

<212> Protein

<213> Artificial Sequence

<220>

<221> misc-feature

<222> Description of Artificial Sequence: Synthetic polypeptide of Core TM

<400> 18

Asp Gln Lys Cys Lys Cys Ala Arg Ile Thr Ser Arg Ile Ile Arg Ser

Ser Glu Asp Pro Asn Glu Asp Ile Val Glu Arg Asn Ile Arg Ile Ile

Val Pro Leu Asn Asn Arg Glu Asn Ile Ser Asp Pro Thr Ser Pro Leu 40

Arg Thr Arg Phe Val Tyr His Leu Ser Asp Leu Cys Lys Lys Asp Glu 55

Asp Ser Ala Thr Glu Thr Cys 70

<210> 19

<211> 49

<212> Protein

<213> Artificial Sequence

<220>

<221> misc-feature

<222> Description of Artificial Sequence: Synthetic polypeptide of C2 fragment

<400> 19

Ser Arg Ile Ile Arg Ser Ser Glu Asp Pro Asn Glu Asp Ile Val Glu

Arg Asn Ile Arg Ile Ile Val Pro Leu Asn Asn Arg Glu Asn Ile Ser 25

Asp Pro Thr Ser Pro Leu Arg Thr Arg Phe Val Tyr His Leu Ser Asp 35 40

Leu

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<210> 20
 <211> 12
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 <213> Artificial Sequence
 <220>
 <221> misc-feature
 <222> Description of Artificial Sequence: Synthetic polypeptide of D 1.1
 fragment
 <400> 20
      Asp Gln Lys Cys Lys Cys Ala Arg Ile Thr Ser Arg
 <210> 21
 <211> 14
 <212> Protein
<213> Artificial Sequence
 <220>
 <221> misc-feature
 <222> Description of Artificial Sequence: Synthetic polypeptide of L3D
 fragment
 <400> 21
 Asp Leu Cys Lys Lys Asp Glu Asp Ser Ala Thr Glu Thr Cys
 <210> 22
 <211> 36
 <212> Protein
 <213> Artificial Sequence
 <220>
 <221> misc-feature
<222> Description of Artificial Sequence: Synthetic polypeptide of T4
fragment
 <400> 22
      Ser Thr Tyr Asp Arg Asn Lys Cys Tyr Thr Ala Val Val Pro Leu Val
                     5
                                         10
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Tyr Gly Glu Thr Lys Met Val Glu Thr Ala Leu Thr Pro Asp Ala

25

20

Cys Tyr Pro Asp 35

33

Protein

Artificial Sequence

₹220>

<221> misc-feature

<222> Description of Artificial Sequence: Synthetic polypeptide of Core TM $c\mathfrak{B}\overline{m}A$ using L3

<400> 23

Asp Asn Lys Cys Lys Cys Ala Arg Ile Thr Ser Arg Ile Ile Arg Ser 1 5 10 15

Ser Glu Asp Pro Asn Glu Asp Ile Val Glu Arg Asn Ile Arg Ile Ile 20 25 30

Val Pro Leu Asn Asn Arg Glu Asn Ile Ser Asp Pro Thr Ser Pro Leu
35 40 45

Arg Thr Arg Phe Val Tyr His Leu Ser Asp Leu Cys Lys Lys Cys Asp 50 55

Pro Thr Glu Val Glu Leu Asp Asn Gln Ile Val Thr Ala Thr Gln Ser 65 70 75 80

Asn Ile Cys Asp Glu Asp Ser Ala Thr Glu Thr Cys Tyr 85

<210> 24

<211> 35

<212> Protein

<213> Artificial Sequence

<220>

<221> misc-feature

<222> Description of Artificial Sequence: Synthetic polypeptide of L3 fragment

<400> 24

Asp Leu Cys Lys Lys Cys Asp Pro Thr Glu Val Glu Leu Asp Asn Gln
1 10 15

Ile Val Thr Ala Thr Gln Ser Asn Ile Cys Asp Glu Asp Ser Ala Thr . 20 25 30

Leu Trp Thr

<210> 25

·-=

<211> 22

<212> Protein

<213> Artificial Sequence

<220>

<221> misc-feature

<222> Description of Artificial Sequence: Synthetic polypeptide of D1 fragment

<400> 25

Asp Glu Asp Glu Arg Ile Val Leu Val Asp Asn Lys Cys Lys Cys 1 5 10 15

Ala Arg Ile Thr Ser Arg

<210> 26

<211> 66

<212> Protein

<213> Artificial Sequence

<220>

<221> misc-feature

<222> Description of Artificial Sequence: Synthetic polypeptide of TpS2

<400> 26

Cys Ser Asp Asp Asp Lys Ala Gln Thr Glu Thr Cys Thr Val Ala 1 5 10 15

Pro Arg Glu Arg Gln Asn Cys Gly Phe Pro Gly Val Thr Pro Ser Gln 20 25 30

Cys Ala Asn Lys Gly Cys Cys Phe Asp Asp Thr Val Arg Gly Val Pro 35 40 45

Trp Cys Phe Tyr Pro Asn Thr Ile Asp Val Pro Pro Glu Glu Glu Cys
50 55

<210> 27

· <211> 421

<212> DNA

<213> Artificial Sequence

<220>

<221> misc-feature

<222> Description of Artificial Sequence: Complementary nucleotide sequence of "full length" TM cDNA

<400> 27

CTAGTCCTTC TACTTGCATA ACAAGACCAA CTGTTGTTCA CGTTCACACG AGCATAATGA

· ic



AGATCTTAGT AGGCATCGAG TCTCCTGGGT TTACTTCTAT ATCAGCTTGC ATTGTAGGCA	120
TAGTAGCAGG GIGACTTATT GGCCCTCTTA TAGAGTCTAG GATGTTCAGG CAACGCGTGT	180
GCGAAGCATA TGGTGGACAG TCTAGACACA TTCTTCACAC TAGGTTGTCT CCATCTCGAC	240
CTGTTAGTCT ATCAGTGACG CTGAGTTTCG TTGTAAACGC TACTCCTGTC GCGATGTCTT	300
TGGACGTCGT GGATGCTATC CTTGTTTACG ATGTGCCGGC ACCAAGGCGA GCACATACCA	360
CCTCTCTGTT TTTACCACCT TTGACGGGAA TGCGGGCTAC GTACGATAGG CCTGACTTAA	.420
G	421
<210> 28 <211> 219 <212> DNA <213> Artificial Sequence	
<pre><221> misc-feature <222> Description of Artificial Sequence: Complementary nucleotide of Core TM cDNA</pre>	s e quence
<400> 28 CTAGTCTTCA CGTTCACACG AGCATAATGA AGATCTTAGT AGGCATCGAG TCTCCTGGGT	60
TTACTTCTAT ATCAGCTTGC ATTGTAGGCA TAGTAGCAGG GTGACTTATT GGCCCTCTTA	120
TAGAGTCTAG GATGTTCAGG CAACGCGTGT GCGAAGCATA TGGTGGACAG TCTAGACACA	180
TTCTTCCTAC TCCTGTCGCG ATGTCTTTGG ACGACTTAA	219
<210> 29 <211> 140 <212> DNA <213> Artificial Sequence	
<220> <221> misc-feature . <222> Description of Artificial Sequence: Complementary nucleotide of C2 fragment	sequ ence
<400> 29 TTAGTAGGCA TCGAGTCTCC TGGGTTTACT TCTATATCAG CTTGCATTGT AGGCATAGTA	60
GCAGGGTGAC TTATTGGCCC TCTTATAGAG TCTAGGATGT TCAGGCAACG CGTGTGCGAA	120

GCATATGGTG GACAGTCTAG

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<210> 30 <211> 31 <212> DNA <220>

<213> Artificial Sequence

<221> misc-feature

<222> Description of Artificial Sequence: Complementary nucleotide sequence of D 1.1 fragment

<400> 30

TCTTCACGTT CACACGAGCA TAATGAAGAT C

31

<210> 31 <211> 44

<212> DNA

<213> Artificial Sequence

<220>

<221> misc-feature

<222> Complementary nucleotide sequence of L3D fragment

<400> 31

ACACATTCTT CCTACTTCTC AGGCGATGTC TTTGGACGAC TTAA

44

<210> 32

<211> 117

<212> DNA

<213> Artificial Sequence

<220>

<221> misc-feature

<222> Description of Artificial Sequence: Complementary nucleotide sequence

of T4 fragment

<400> 32

ACGTCGTGGA TGCTATCCTT GTTTACGATG TGCCGGCACC AAGGCGAGCA CATACCACCT 60

CTCTGTTTTT ACCACCTTTG ACGGGAATGC GGGCTACGTA CGATGGGACT GACTTAA 117

-12



<210> 33	•	
<211> 282		
<212> DNA	•	
<213> Artificial Sequence		•
in annual podecies		
<220>		
<221> misc-feature		
<222> Description of Artificial	Sequence: Complementary nucleotide	sednence
of Core TM cDNA using L3		
<400> 33		
CTGTTGTTCA CGTTCACACG AGCATAATGA	AGATCTTAGT AGGCATCGAG TCTCCTGGGT	6 0
		60
ምዋልሮፓምር ዋልዋ ል ጥሮል ਫሮንምਫሮ ልጥነሩምልਫਫሮል	TAGTAGCAGG GTGACTTATT GGCCCTCTTA	120
- WINDERSTON METALOGOM	TAGIRGUAG GIGACTIRIT GGCCCICTIA	120
Тасастство свтетсвое свлосостен	GCGAAGCATA TGGTGGACAG TCTAGACACA	
INDICING GRIGITERGG CAMCGCGIGI	GCGAAGCAIA IGGIGGACAG TCTAGACACA	180
TECTECICIO EN COMPOSEDE CON MORGO O		
TTCTTCACAC TAGGTTGTCT CCATCTCGAC	CTGTTAGTCT ATCAGTGACG CTGAGTTTCG	240
EMCC133466		
TTGTAAACGC TACTCCTGTC GCGATGTCTT	TGGACGATGA CT	282
		•
•		
<210> 34		
<211> 105		
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<213> Artificial Sequence		
		•
<220>		
<221> misc-feature		
<222> Description of Artificial	Sequence: Complementary nucleotide	sequence
of L3 fragment		•
	•	
<400> 34		
GATCTGTGTA AGAAGTGTGA TCCAACAGAG	GTAGAGCTGG ACAATCAGAT AGTCACTGCG	60
		0.0
ACTCAAAGCA ACATTTGCGA TGAGGACAGC	GCTACACTTT GGACG	105
		105
<210> 35		
<211> 65		
<212> DNA		
<213> Artificial Sequence		
versa sednence		
<220>		
<221> misc-feature .		
	Communication of the contract of the	
es Dr emanage of Artificial	Sequence: Complementary nucleotide	Redneuce
of D1 fragment	•	
-100: 25		
<400> 35		
CTAGTCCTTC TACTTGCATA ACAAGACCAA	CTGTTGTTCA CGTTCACACG AGCATAATGA	60
3 C2 MA		<u>.</u>
AGATC		65

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<210> 36
  <211> 206
  <212> DNA
  <213> Artificial Sequence
  <220>
  <221> misc-feature
  <222> Description of Artificial Sequence: Complementary nucleotide sequence
  <400> 36
 ACTTCGCTAC TGCTGCTATT CCGGGTTTGC CTCTGGACAT GACAACGCGG AGCACTTGCC
                                                                        60
 GTTTTGACGC CTAAGGGCCT TCATTGTGGG AGAGTCACGC GATTATTTCC GACGACAAAA
                                                                       120
 CTACTGTGCC ATGCCCCGCA AGGCACCACG AAGATGGGGT TATGTTAACT GCAAGGGGGA
                                                                       180
 CTTCTTCTCA CGCTCGGCAT TCTTAA
                                                                       206
  <210> 37
  <211> 13
  <212> Protein
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 <220>
 <221> misc-feature
 <2222> Description of Artificial Sequence: Domain 1, 13 amino peptide with
 substantial β-sheet character
· <400>. 37
      Asp Gln Glu Asp Glu Arg Ile Val Leu Val Asp Asn Lys
                      5
 <210> 38
 <211> 7
 <212> Protein
 <213> Tobacco etch virus
 <220>
 <221> misc-feature
 <222> Peptide recognized by the tobacco etch virus protease Nia
 <400> 38
      Glu Asn Leu Tyr Phe Gln Ser
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<210> 39
<211> 11
<212> Protein
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<222> Description of Artificial Sequence: Synthetic polypeptide residues
from pro-cathepsin E
<400> 39
    Lys Ala His Lys Val Asp Met Val Gln Tyr Thr
<210> 40
<211> 4
<212> Protein
<213> Artificial Sequence
<220>
<221> misc-feature
<222> Description of Artificial Sequence: Linker from procathepsin
<400> 40
    Val Gln Tyr Thr
   . 1
<210> 41
<211> 6
<212> Protein
<213> Human
<220>
<221> misc-feature
<222> Linker from polyimmunoglobulin receptor
<400> 41
    Glu Lys Ala Val Ala Asp
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<210> 42 <211> 131 <212> DNa <213> Artificial Sequence <220> CDS <221> 1..78 <222> Description of Artificial Sequence; Nucleotide sequence of secretion signal from pMelBac <400> 42 ATG AAA TTC TTA GTC AAC GTT GCC CTT TTT ATG GTC GTA TAC ATT TCT 48 Met Lys Phe Leu Val Asn Val Ala Leu Phe Met Val Val Tyr Ile Ser 40 TAC ATC TAT GCG GAT CCG AGC TCG AGT GCT CTAGATCTGC AGCTGGTACC 98 Tyr Ile Tyr Ala Asp Pro Ser Ser Ser Ala 55 - ATGGAATTCG AAGCTTGGAG TCGACTCTGC TGA 131 <210> 43 <211> 26 <212> Protein <213> Artificial Sequence <220> <221> misc-feature <222> Description of Artificial Sequence: Synthetic polypeptide sequence of secretion signal from pMelBac <400> 43 Met Lys Phe Leu Val Asn Val Ala Leu Phe Met Val Val Tyr Ile Ser 1 5 Tyr Ile Tyr Ala Asp Pro Ser Ser Ser Ala 20 <210> 44 <211> 4 <212> Protein <213> Artificial Sequence <220>

<222> Description of Artificial Sequence: Endomembrane retention signal

<221> misc-feature

Lys Asp Glu Leu

<400> 44

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<210> 45
 <211> 16
 <212> Protein
 <213> Human
 <220>
 <221> misc-feature
 <222> Residues 585-600 of polyimmunoglobulin receptor
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      Ala Ile Gln Asp Pro Arg Leu Phe Ala Glu Glu Lys Ala Val Ala Asp
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                                                                       61
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 CTAGAAGTAA TACGAGCACA CTTGCACTTG TTGTCAACCA GAACAATACG TTCATCTTCC
                                                                       60
T
                                                                       61
<210> 48
 <211> 31
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 <221> misc-feature
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 GATCAGAAGT GCAAGTGTGC TCGTATTACT T
                                                                       31
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60

61

<222> Description of Artificial Sequence: Oligonucleotide 2.2ser

CTAGAAGTAA TACGAGCGGA CTTGCACTTG TTGTCAACCA GAACAATACG TTCATCTTCC

<400> 51

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<210> 52
<211> 61
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                                                                      61
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CTAGAATCAT CCGTAGCTCA GAGGACCCAA ATGAAGATAT AGTCGAA
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GATACGGATG TTACGTTCGA CTATATCTTC ATTTGGGTCC TCTGAGCTAC GGATGATT
                                                                      58
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49

ACGGACTTGT AGGATCTGAG ATGTGCTCCC GGTTATTCAG TGGGACGAT

44

GATCTGTGTA AGAAGGATGA GGACAGCGCT ACAGAAACCT GCTG

<400> 63

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59

GATCTGTGTA AGAAGTCTGA TATCGATGAA GATTCCGCTA CAGAAACCTG CAGCACATG

<400> 67

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	, ° ° ' ' ' ' ' ' ' ' ' ' ' ' ' ' ' ' '	
<212>		
	Dela Dela Control Cont	
ĕŘ₹RA.	Artificial Sequence	
<220>		
	nian fantum	
	misc-feature	
<222>	Description of Artificial Sequence: Oligonucleotide 10.2A3	
<400>	69	
	PGTG CTGCAGGTTT CTGTAGCGGA ATCTTCATCG ATATCAGACT TCTTACACA	59
MATICA.	TOTO CIGCAGOTTI CIGIROCGGA ATCITCATCG ATAICAGACT TCTIACACA	22
	·	
<210>	69	
<211>	64	
<212>	DNA	
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CTAA		54
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<222> Description of Artificial Sequence: Oligonucleotide 10.3\(\Delta\)/ser68

<400> 70

AATCTTCATC GATATCAGAC TTCTTAGACA

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s <u>s</u>

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<210> 71
<211> 64
<212> DNA
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CTAA
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ATTGTCCAGC TCTACCTCTG TTGGATCACA CTTCTTACAC A
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<210> 74
<211> 46
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<221> misc-feature
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<400> 74
                                                                      46
ACTCAAAGCA ACATTTGCGA TGAGGACAGC GCTACAGAAA CCTGCA
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50

<211> 50
<212> DNA
<213> Artificial Sequence
<220>
<221> misc-feature
<222> Description of Artificial Sequence: Oligonucleotide 15
<400> 78
ACAAAAATGG TGGAAACTGC CCTTACGCCC GATGCATGCT ATCCGGACTG

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<210>	79	
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<212>	DNA	
<213>	Artificial Sequence	
	•	
<220>		
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	Description of Artificial Sequence: Oligonucleotide 16	
44447	Descripcion of Michaelar and arms. Arms.	
<400>		60
AATTCA	GTCC GGATAGCATG CATCGGGCGT AAGGGCAGTT TCCACCATTT TTGTCTCTCC	90
ACCATA	CAC	69
<210>	80	
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<213>	Artificial Sequence	
	•	
<220>		
	misc-feature	
	Description of Artificial Sequence: Oligonucleotide 15KDEL	
-4242	DDDGTTBatam de tweethert pudgement estalementes tormes	
<400>	80	
•	AATGG TGGAAACTGC CCTTACGCCC GATGCATGCT ATCCGGACAA GGATGAATTG	60
NCARA	Wide Idawarcide chiireace ariachiaci Riccadacar annina	UV
TĠ		62
16		02
014	**	
<210>		
<211>	- -	
<212>		
<213>	Artificial Sequence	
<220>		
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<222>	Description of Artificial Sequence: Oligonucleotide 16KDBL	
<400>	81	
	ACANT TOATCOTTOT COGGATAGON TGCATOGGGO GTANGGGOAG TTTCCACCAT	60
وكالماسليك	TCTCT CCACCATACA C	81
		~ -

421U2	82	
<211>	88	
<212>	DNA	
<213>	Artificial Sequence	
<220>		
	misc-feature	
<222>	Description of Artificial Sequence: Oligonucleotide Pl	
<400>	82	
GATCAG	GTCG CTGCCATCCA AGACCCGAGG CTGTTCGCCG AAGAGAAGGC CGTCGCTGAC	60
		•
ጥሮሮኒኒኒኒኒኒኒኒኒኒኒኒኒኒኒኒኒኒኒኒኒኒኒኒኒኒኒኒኒኒኒኒኒኒኒኒ	TGCA AGTGTGCTCG TATTACTT	88
ICCNAO	IOCA ADIGIOCICO IMITACII	86
<210>	•	
<211>	88	
<212>	DNA	
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	-	
<220>		
	misc-feature	
42222	Description of Artificial Sequence: Oligonucleotide P2	
	,	
<400>		
CTAGAA	GTAA TACGAGCACA CTTGCACTTG GAGTCAGCGA CGGCCTTCTC TTCGGCGAAC	60
AGCCTC	GGGT CTTGGATGGC AGCGACCT	8 B
<210>	84	
<211>	10	
	Protein	
	Artificial Sequence	
(2132	Arcittotat Bequence	
•••	•	
<220>		
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<400>	84	
Ċ	ys Ala Ala Pro Lys Lys Arg Lys Val	
1		
-	3 40	

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<210> 85
<211> 22
<212> Protein
<213> Artificial Sequence
<220>
<221> misc-feature
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     Cys Ala Ala Lys Arg Pro Pro Ala Ala Ile Lys Lys Ala Ala Ala Gly
                     5
                                        10
     Gln Ala Lys Lys Lys Lys
<210> 86
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intracellular targeting
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    His Asp Glu Leu
<210> 67
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GCGATGACGA CGATAAGGCC CAAACGGAGA CCTGTACTGT TGCGCCTCGT GAACGGCAAA
                                                                      60
ACTGCGGATT CCCGGAA
                                                                      77
```

<210>	.88		
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	Artificial Sequence		
~	wetrrer oedrence		
<220>			
	misc-feature	•	
<222>	-	Sequence: Oligonucleotide Tp2	
~~~~	Descripcion of Attiticial	pedrence: offBouncieocrae ibs	
<400>	98		
41002			
GTTTTG	ACATBACAAC DOBDABCACT TECC	GGTCTCCGTT TGGGCCTTAT CGTCGTCATC 6	0
		designed a second secon	•
GCTTCA		6	6
		•	_
<210>	89		
<211>	72		
<212>	DNA		
<213>	Artificial Sequence		
	<u>-</u>		
<220>			
<221>	misc-feature		
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		-	
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GTAACA	CCT CTCAGTGCGC TAATAAAGGC	TGCTGTTTTG ATGACACGGT ACGGGGCGTT 6	O
•			
CCGTGG	rgct tc	7	2
<210>			
<211>			
<212>			
<213>	Artificial Sequence		
<220>			
	misc-feature	A 1	
<2225	Description of Arthricial	Sequence: Oligonucleotide Tp4	
<400>	90		
		TTTATTAGCG CACTGAGAGG GTGTTACTTC 6	0
GCCCCG.	HACE GIGICALLAN MACAGCAGCC	IIIAIIAGG CACIGAGAGG GIGIIACIIC 6	U
CGGGAA	rccg ca	7	2
~~~~~	VFF	•	-

₹210>	91	
<211>	49	
<212>	DNA	
<213>	Artificial Sequence	
	•	
<220>	•	
<221>	misc-feature	
<222>	Description of Artificial Sequence: Oligonucleotide Tp5	
1	• • • • • • • • • • • • • • • • • • •	
<400>	91	
TACCO	CAATA CAATTGACGT TCCGCCTGAA GAAGAGTGCG AGCCGTAAG	49
	· · · · · · · · · · · · · · · · · · ·	•
	•	
<210>	92	
<211>	68	
<212>	DNA	
<213>	Artificial Sequence	
<220>		
<221>	misc-feature	
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CACGG	AAC .	68
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<222>	Description of Artificial Sequence: Synthetic peptide linker	
	• • • • • • • • • • • • • • • • • • • •	
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	1 5 10	